In the claims:

Amend claims 8, 28, and 29 as follows:

--8. (Amended) The peptide of claim 6, wherein the amino acid sequence is selected from the group consisting of Ala-Gln-Lys-Arg-Arg (SEQ ID NO:124), Gly-Lys-Ser-Arg-Arg (SEQ ID NO:125), Glu-Gln-Lys-Arg-Arg (SEQ ID NO:126), Glu-Ala-Lys-Arg-Arg (SEQ ID NO:127), Gly-Gln-Lys-Arg-Arg (SEQ ID NO:128), Gly-Ala-Lys-Arg-Arg (SEQ ID NO:129), Gly-Lys-Lys-Arg-Arg (SEQ ID NO:130), Gly-His-Lys-Arg-Arg (SEQ ID NO:131), Gly-Lys-Ala-Phe-Arg (SEQ ID NO:132), Glu-Lys-Ala-Gln-Arg (SEQ ID NO:133), and Glu-Lys-Ala-Arg-Arg (SEQ ID NO:134).--

--28. (Amended) The composition of claim 17, wherein the peptide is Gly-Gly-Lys-Ala-Arg-Arg-Leu (SEQ ID NO:135).--

--29. (Amended) The composition of claim 17, wherein the therapeutic drug is a compound belonging to the group of thapsigargins which have been derivatized with a moiety containing a primary amine group, the peptide is Gly-Gly-Lys-Ala-Arg-Arg-Leu (SEQ ID NO:135), and the linker is selected from the group consisting of unsubstituted or alkyl-, aryl-, halo-, alkoxy-, alkenyl-, amido- or amino-substituted CO-$(CH=CH)_{n1}$-$(CH_2)_{n2}$-Ar-NH_2, CO-$(CH_2)_{n2}$-$(CH=CH)_{n1}$-Ar-NH_2, CO-$(CH_2)_{n2}$-$(CH=CH)_{n1}$-CO-NH-Ar-NH_2, CO-$(CH=CH)_{n1}$-$(CH_2)_{n2}$-CO-NH-Ar-NH_2, CO-$(CH_2)_{n3}$-NH_2, and CO-$(CH_2)_{n3}$-NH-CO-CH(R_4)-NH_2, wherein n1 and n2 are from 0 to 5, n3 is from 0 to 15, Ar is any substituted or unsubstituted aryl group, attachment of NH_2 to Ar is in a ortho, meta or para position with respect to the remainder of the linker, and R_4 is any naturally occurring amino acid side chain.--

In the claims:

Claims 8, 28, and 29 have been amended as follows:

8. (Amended) The peptide of claim 6, wherein the amino acid sequence is selected from the group consisting of Ala-Gln-Lys-Arg-Arg (SEQ ID NO:124), Gly-Lys-Ser-Arg-Arg (SEQ ID NO:125), Glu-Gln-Lys-Arg-Arg (SEQ ID NO:126), Glu-Ala-Lys-Arg-Arg (SEQ ID NO:127), Gly-Gln-Lys-Arg-Arg (SEQ ID NO:128), Gly-Ala-Lys-Arg-Arg (SEQ ID NO:129), Gly-Lys-Lys-Arg-Arg (SEQ ID NO:130), Gly-His-Lys-Arg-Arg (SEQ ID NO:131), Gly-Lys-Ala-Phe-Arg (SEQ ID NO:132), Glu-Lys-Ala-Gln-Arg (SEQ ID NO:133), and Glu-Lys-Ala-Arg-Arg (SEQ ID NO:134).

28. (Amended) The composition of claim 17, wherein the peptide is Gly-Gly-Lys-Ala-Arg-Arg-Leu (SEQ ID NO:135).

29. (Amended) The composition of claim 17, wherein the therapeutic drug is a compound belonging to the group of thapsigargins which have been derivatized with a moiety containing a primary amine group, the peptide is Gly-Gly-Lys-Ala-Arg-Arg-Leu (SEQ ID NO:135), and the linker is selected from the group consisting of unsubstituted or alkyl-, aryl-, halo-, alkoxy-, alkenyl-, amido- or amino-substituted $CO\text{-}(CH{=}CH)_{n1}\text{-}(CH_2)_{n2}\text{-}Ar\text{-}NH_2$, $CO\text{-}(CH_2)_{n2}\text{-}(CH{=}CH)_{n1}\text{-}Ar\text{-}NH_2$, $CO\text{-}(CH_2)_{n2}\text{-}(CH{=}CH)_{n1}\text{-}CO\text{-}NH\text{-}Ar\text{-}NH_2$, $CO\text{-}(CH{=}CH)_{n1}\text{-}(CH_2)_{n2}\text{-}CO\text{-}NH\text{-}Ar\text{-}NH_2$, $CO\text{-}(CH_2)_{n3}\text{-}NH_2$, and $CO\text{-}(CH_2)_{n3}\text{-}NH\text{-}CO\text{-}CH(R_4)\text{-}NH_2$, wherein n1 and n2 are from 0 to 5, n3 is from 0 to 15, Ar is any substituted or unsubstituted aryl group, attachment of NH_2 to Ar is in a ortho, meta or para position with respect to the remainder of the linker, and R_4 is any naturally occurring amino acid side chain.